

Serial Number: 09/955807

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data" **ENTERED**
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☒ Other: Format for applicant was incorrect. Erroneous field identifier inserted within text. mt

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

#2

OIPE

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/955,807

DATE: 10/16/2001

TIME: 09:42:56

Input Set : N:\Crf3\10042001\I955807.raw

Output Set: N:\CRF3\10162001\I955807.raw

ENTERED

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1 <110> APPLICANT: Lok, Si
2      Sheppard, Paul O.
3      Kindsvogel, Wayne
4      Bort, Susan J.
5 <120> TITLE OF INVENTION: Secretory Protein-48
6 <130> FILE REFERENCE: 98-17C1
7 <140> CURRENT APPLICATION NUMBER: US/09/955,807
8 <141> CURRENT FILING DATE: 2001-09-19
9 <150> PRIOR APPLICATION NUMBER: 60/102,679
10 <151> PRIOR FILING DATE: 1998-10-01
11 <150> PRIOR APPLICATION NUMBER: 09/410,603
12 <151> PRIOR FILING DATE: 1999-10-01
13 <160> NUMBER OF SEQ ID NOS: 17
14 <170> SOFTWARE: FastSEQ for Windows Version 3.0
16 <210> SEQ ID NO: 1
17 <211> LENGTH: 1692
18 <212> TYPE: DNA
19 <213> ORGANISM: Homo sapiens
20 <220> FEATURE:
21 <221> NAME/KEY: CDS
22 <222> LOCATION: (59)...(373)
23 <400> SEQUENCE: 1
24      tttttctaag ggatgagata agaataaata gaaatttttg catttcttct cacattag      58
25      atg ctg ggt tat tct gag ccc atg cca tgt gca cac cca ctt ggc ctc      106
26      Met Leu Gly Tyr Ser Glu Pro Met Pro Cys Ala His Pro Leu Gly Leu
27      1          5          10          15
28      ttc ctc tta ggc cta cac cct gcc ctt tct ttg ccc ctt gta gtt act      154
29      Phe Leu Leu Gly Leu His Pro Ala Leu Ser Leu Pro Leu Val Val Thr
30      20          25          30
31      gtg gct gga gtg atg agc gcc act ccc aag cat ggc ctg gaa caa tgt      202
32      Val Ala Gly Val Met Ser Ala Thr Pro Lys His Gly Leu Glu Gln Cys
33      35          40          45
34      cct cct gcc cct cca cca gca gtg aca gga ttc act ggg gac tcg ggg      250
35      Pro Pro Ala Pro Pro Pro Ala Val Thr Gly Phe Thr Gly Asp Ser Gly
36      50          55          60
37      gca aag gag act gtg tca caa gac aaa agg agc cag ggt cac aca tgg      298
38      Ala Lys Glu Thr Val Ser Gln Asp Lys Arg Ser Gln Gly His Thr Trp
39      65          70          75          80
40      tgt acc ctc gcc ctg cct cac cca tgg ctg aca tgg gtt gga cac ctc      346
41      Cys Thr Leu Ala Leu Pro His Pro Trp Leu Thr Trp Val Gly His Leu
42      85          90          95
43      aga aat cat gtg tct tca gcg agc cac tgagagttgg ggcttttatct      393
44      Arg Asn His Val Ser Ser Ala Ser His
45      100          105
46      gttactcggc taggggtaac ctaaccgatg agactgtaac tggttactgt aaataaccaa      453
47      gctcccagta atagtaaacc agtgacaaaa acaattctta tccaaaaagg ttcacctttt      513
48      tttaaaatgt gtgaactaaa acagtcttta ttgctctaag acattaaaat ttgcactttt      573

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DATE: 10/16/2001

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Output Set: N:\CRF3\10162001\I955807.raw

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49      ttgatgttga ataccactga atatttttatt tttatatattt attacacaga aatacagcaa      633
50      ttattacaaa acgagtatta ggaatggcaa aggctttagg acagactatt agcggaaaac      693
51      atttggaact taaggagtgt tttacatttg gaacttactt taaggagtgt cgttcagaca      753
52      ctactatat cttaacctca gtttttagaa gtaagcaagc tctcattttt tgctattcat      813
53      atttgaagtg attaaactca taaatttgaa atttactttt tagagaccaa agattaaaat      873
54      taggtgggat gtcagctttt aaaatatact aagatttcct acaactacca atagcttatt      933
55      tccctgggaa acagattaca ttgtagtact taaccagaa ctcatgcagt tcatccaaaa      993
56      tgatggtaaa cttttttcct cagaattacc taactttcct tgactatgaa ttcaacattc     1053
57      aagaatcttc ttctggtagc aggagcggca gagaggacag gcatggaaaag gaggcctgtc     1113
58      tcccacggag aactcctcta gtgccagcag acacgcatgg tggaacacat gtgagcagga     1173
59      caggagggcc atctctctgg aacgcctgcc cgcaccacg cactgaccgc cagcagcgga     1233
60      gagaggggac aggcagatgg agcactcctg ggtctcccg cgcagagact gcggcacaca     1293
61      ggacaggaag aggccacgcg ggttagtttc atcacagcag aaagttaactt aaactgaaat     1353
62      gcgaaccatg tgccccgaga catgggtctt cgaaacatgc ggaagtttca ttctgtgtta     1413
63      aaatcacatg cattttattt atatatatac atatatatat atacacacac acatatactc     1473
64      tgttactcct gggaactgtg gaaaggggta gtaaccacac tgtgataagc aacatccaac     1533
65      aggaacttcc agaatttcaa actgaaggga cctttgccgt caccctaaag cccatgagga     1593
66      aagtcctacc acaggtgcag gggcagctag ggcagcggtt accccaggcc tgacactcct     1653
67      aggcttccca aagtgaagtcc tcgacctccc ccgctcgag                               1692

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69 &lt;210&gt; SEQ ID NO: 2

70 &lt;211&gt; LENGTH: 105

71 &lt;212&gt; TYPE: PRT

72 &lt;213&gt; ORGANISM: Homo sapiens

73 &lt;400&gt; SEQUENCE: 2

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74      Met Leu Gly Tyr Ser Glu Pro Met Pro Cys Ala His Pro Leu Gly Leu
75      1          5          10          15
76      Phe Leu Leu Gly Leu His Pro Ala Leu Ser Leu Pro Leu Val Val Thr
77      20          25          30
78      Val Ala Gly Val Met Ser Ala Thr Pro Lys His Gly Leu Glu Gln Cys
79      35          40          45
80      Pro Pro Ala Pro Pro Pro Ala Val Thr Gly Phe Thr Gly Asp Ser Gly
81      50          55          60
82      Ala Lys Glu Thr Val Ser Gln Asp Lys Arg Ser Gln Gly His Thr Trp
83      65          70          75          80
84      Cys Thr Leu Ala Leu Pro His Pro Trp Leu Thr Trp Val Gly His Leu
85      85          90          95
86      Arg Asn His Val Ser Ser Ala Ser His
87      100          105

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89 &lt;210&gt; SEQ ID NO: 3

90 &lt;211&gt; LENGTH: 79

91 &lt;212&gt; TYPE: PRT

92 &lt;213&gt; ORGANISM: Homo sapiens

93 &lt;400&gt; SEQUENCE: 3

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94      Leu Pro Leu Val Val Thr Val Ala Gly Val Met Ser Ala Thr Pro Lys
95      1          5          10          15
96      His Gly Leu Glu Gln Cys Pro Pro Ala Pro Pro Pro Ala Val Thr Gly
97      20          25          30
98      Phe Thr Gly Asp Ser Gly Ala Lys Glu Thr Val Ser Gln Asp Lys Arg
99      35          40          45

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100      Ser Gln Gly His Thr Trp Cys Thr Leu Ala Leu Pro His Pro Trp Leu
101          50                      55                      60
102      Thr Trp Val Gly His Leu Arg Asn His Val Ser Ser Ala Ser His
103          65                      70                      75
105 <210> SEQ ID NO: 4
106 <211> LENGTH: 77
107 <212> TYPE: PRT
108 <213> ORGANISM: Homo sapiens
109 <400> SEQUENCE: 4
110      Leu Val Val Thr Val Ala Gly Val Met Ser Ala Thr Pro Lys His Gly
111          1                      5                      10                      15
112      Leu Glu Gln Cys Pro Pro Ala Pro Pro Ala Val Thr Gly Phe Thr
113          20                      25                      30
114      Gly Asp Ser Gly Ala Lys Glu Thr Val Ser Gln Asp Lys Arg Ser Gln
115          35                      40                      45
116      Gly His Thr Trp Cys Thr Leu Ala Leu Pro His Pro Trp Leu Thr Trp
117          50                      55                      60
118      Val Gly His Leu Arg Asn His Val Ser Ser Ala Ser His
119          65                      70                      75
121 <210> SEQ ID NO: 5
122 <211> LENGTH: 65
123 <212> TYPE: PRT
124 <213> ORGANISM: Homo sapiens
125 <400> SEQUENCE: 5
126      Pro Lys His Gly Leu Glu Gln Cys Pro Pro Ala Pro Pro Pro Ala Val
127          1                      5                      10                      15
128      Thr Gly Phe Thr Gly Asp Ser Gly Ala Lys Glu Thr Val Ser Gln Asp
129          20                      25                      30
130      Lys Arg Ser Gln Gly His Thr Trp Cys Thr Leu Ala Leu Pro His Pro
131          35                      40                      45
132      Trp Leu Thr Trp Val Gly His Leu Arg Asn His Val Ser Ser Ala Ser
133          50                      55                      60
134      His
135          65
137 <210> SEQ ID NO: 6
138 <211> LENGTH: 384
139 <212> TYPE: DNA
140 <213> ORGANISM: Homo sapiens
141 <220> FEATURE:
142 <221> NAME/KEY: variation
143 <222> LOCATION: (1)...(384)
144 <223> OTHER INFORMATION: n is any nucleotide
145 <221> NAME/KEY: misc_feature
146 <222> LOCATION: (1)...(384)
147 <223> OTHER INFORMATION: n = A,T,C or G ✓
148 <400> SEQUENCE: 6
149      tttttctaag g gatgagata agaataaata gaaattttgg catttcttct cacattagat      60
150      gctgggttat tctgagccca tgccatgtgc acaccactt ggcctcttcc tcttaggcct      120
151      acacctgcc ctttctttgc ccctttagt tactgtggct ggagtgatga ggcacctcc      180

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TIME: 09:42:56

Input Set : N:\Cr3\10042001\I955807.raw

Output Set: N:\CRF3\10162001\I955807.raw

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152      caagcatggc ctggaacaat gtcctcctgc ccctccacca gcagtgcacag gattcactgg      240
153      ggactcgggg gcaaaggaga ctgtgtcaca agacaaaagg agccagggtc acacatgggtg      300
W--> 154      ttacctcgcc ctgcctcacc catgggtgac atgggttgga cacctcanaa atcntgtttc      360
W--> 155      ttcaccganc cactgaaaat tggg      384
157 <210> SEQ ID NO: 7
158 <211> LENGTH: 48
159 <212> TYPE: DNA
160 <213> ORGANISM: Homo sapiens
161 <400> SEQUENCE: 7
162      gtctgggttc gctactcgag gcggccgcta tttttttttt tttttttt      48
164 <210> SEQ ID NO: 8
165 <211> LENGTH: 20
166 <212> TYPE: PRT
167 <213> ORGANISM: Homo sapiens
168 <400> SEQUENCE: 8
169      Ser Ala Thr Pro Lys His Gly Leu Glu Gln Cys Pro Pro Ala Pro Pro
170      1          5          10          15
171      Pro Ala Val Thr
172      20
174 <210> SEQ ID NO: 9
175 <211> LENGTH: 42
176 <212> TYPE: PRT
177 <213> ORGANISM: Homo sapiens
178 <400> SEQUENCE: 9
179      Ser Ala Thr Pro Lys His Gly Leu Glu Gln Cys Pro Pro Ala Pro Pro
180      1          5          10          15
181      Pro Ala Val Thr Gly Phe Thr Gly Asp Ser Gly Ala Lys Glu Thr Val
182      20          25          30
183      Ser Gln Asp Lys Arg Ser Gln Gly His Thr
184      35          40
186 <210> SEQ ID NO: 10
187 <211> LENGTH: 65
188 <212> TYPE: PRT
189 <213> ORGANISM: Homo sapiens
190 <400> SEQUENCE: 10
191      Ser Ala Thr Pro Lys His Gly Leu Glu Gln Cys Pro Pro Ala Pro Pro
192      1          5          10          15
193      Pro Ala Val Thr Gly Phe Thr Gly Asp Ser Gly Ala Lys Glu Thr Val
194      20          25          30
195      Ser Gln Asp Lys Arg Ser Gln Gly His Thr Trp Cys Thr Leu Ala Leu
196      35          40          45
197      Pro His Pro Trp Leu Thr Trp Val Gly His Leu Arg Asn His Val Ser
198      50          55          60
199      Ser
200      65
202 <210> SEQ ID NO: 11
203 <211> LENGTH: 20
204 <212> TYPE: PRT
205 <213> ORGANISM: Homo sapiens

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Input Set : N:\Crf3\10042001\I955807.raw

Output Set: N:\CRF3\10162001\I955807.raw

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208       1             5             10             15
209   Gln Gly His Thr
210               20
212 <210> SEQ ID NO: 12
213 <211> LENGTH: 43
214 <212> TYPE: PRT
215 <213> ORGANISM: Homo sapiens
216 <400> SEQUENCE: 12
217   Thr Gly Asp Ser Gly Ala Lys Glu Thr Val Ser Gln Asp Lys Arg Ser
218       1             5             10             15
219   Gln Gly His Thr Trp Cys Thr Leu Ala Leu Pro His Pro Trp Leu Thr
220               20             25             30
221   Trp Val Gly His Leu Arg Asn His Val Ser Ser
222       35             40
224 <210> SEQ ID NO: 13
225 <211> LENGTH: 12001
226 <212> TYPE: DNA
227 <213> ORGANISM: Homo sapiens
228 <220> FEATURE:
229 <221> NAME/KEY: CDS
230 <222> LOCATION: (10258)...(10572)
231 <400> SEQUENCE: 13
232   actttgtgca aaattccttc attaacgctt agcttcctta tctgtaatac agtgatagta      60
233   tcatcttcct gttagggttt ttgtgaagat caacggaaat aattctgtaa gatccttagc      120
234   atagcgcttg gcacatccta agaactcagt aaatatttagc ccctttatta tgacgatggt      180
235   ggtcatgggt gtggtgagga tgatacgggt tgaaaagctt atctcttggt aataatacct      240
236   tttagttaaa gcttttttga ggcttggtatt ttgcaagtat taggctaacc cataagtctc      300
237   ttcatthaagc cagagaataa attcaagatg aaaacgttag cattcttggc attgatgtaa      360
238   tagaagagag gggatttact gttatgtgtt ccaagagtca catgtattgt aatgggtgta      420
239   aaaacgggta ggttttagcta aagggtacaa acgtaacctt tgaatgtatt tttatgctta      480
240   tttccacatt agtgctaaac atatttcaag ttttatactt taaaaatacc aggacaaagt      540
241   aaattatctt ggtttggggg gggagggggg tgtaatttta tgacagaaga agggaaaggc      600
242   agtgacttct tgtagaaaat ttttaaaaat cctgacatta gctcatttac ctgagttgac      660
243   atgatttgaa tgcataatgac tccatactgg ggcttttagc tattgtaaaa ggccacatac      720
244   tgatggattc attaaggtcc agttttcaga taacttaaac gatatgagca gcaataaagc      780
245   ttctcagatc accaggcctt tccaaccttg atgtttgaga gggtgacctt tgggaggcac      840
246   aaaagatttc agatgagctg tccatatgta ttttactttg aatatgccct gggaggggat      900
247   ggctcatcaa atattgcaat gcctgacagg aaaaagtcac agctcatttc agctgacaca      960
248   ccagataact tatacctttt aatgcttagg ttaataaaag ctggcccaac ttgaagtagg      1020
249   aatcaaacag tcctttttat cagatgtcta gcattaaaac ttaattttta agcctgttat      1080
250   aatatcagca agattagtta gccatggttt cagataaatt tccactttcc attcgctaaa      1140
251   tgagatgggt gcaaataaac tgccgtaact ttagcttttg aattaggtat tctggacatc      1200
252   attttgctaa gaaagccttt attaaagtaa taaaacataa cctgatataa aaggccttat      1260
253   atgcatgtca gttccttgac cataagagag agtagaatta gcaagagttg tataaaaacta      1320
254   cctaatagat acatttactt ttcttcccca gtgtttttca gtattctttg ggggtgtgcta      1380
255   cggggcaatt tatacataga aaaagagtct tattaagtat atgtaatggt tgaatgatct      1440
256   gagatcttaa cagggattta gctgagacct gtaatttgat tgtaaagtag ctatcccctt      1500

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/955,807

DATE: 10/16/2001

TIME: 09:42:57

Input Set : N:\Crf3\10042001\I955807.raw

Output Set: N:\CRF3\10162001\I955807.raw

L:154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:155 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6